Please replace the paragraph at page 56, lines 10-14 with the replacement paragraph as follows:

Amino acid sequences surrounding the activating phosphorylation sites on PKB, p70 S6 kinase and PKCζ and their similarity to the corresponding regions of SGK. The proposed consensus sequence is shown (SEQ ID NO:45). Identities are shown in boldface type and the phosphorylated residues in PKB and p70 S6 kinase are underlined. The phosphorylation sites are separated by 160-165 residues in each enzyme.

Please replace the paragraph at page 60, line 23 to line 29 with the replacement paragraph as follows:

Fig. 11. Nucleotide and deduced amino acid sequences of human and mouse SGK2. A, SGK2α (SEQ ID NO:7)(Corresponding amino acid sequence shown as SEQ ID NO:1). Residues 33-239 correspond to the kinase catalytic domain. The termination codon is marked by a solid triangle. B, SGK2β; (SEQ ID NO:5)(Corresponding amino acid sequence shown as SEQ ID NO:8); The 5' region of SGK2β where it differs from SGK2α. The initiating methionine which starts the sequence of SGK2α is marked by an asterisk. After this residue, the sequences of SGK2α and SGK2β are identical.

Please replace the paragraph at page 61, line 1 to line 6 with the replacement paragraph as follows:

Fig. 12 Nucleotide (SEQ ID NO: 6) and deduced amino acid sequences (SEQ ID NO:4) of human SGK3. Residues 93-389 correspond to the kinase catalytic domain. The termination codon is marked by a sold triangle, the three asterisks denote the position of the most 5' ATG codon and the termination

codon that immediately precedes it is underlined.

Please replace the paragraph at 61, line 8 to line 14 with the replacement paragraph as follows:

Fig. 13 Alignment of amino acid sequences of SGK isoforms. The alignment of human (h) SGK1, hSGK2 (SEQ ID NO:8), murine (m) SGK2 (SEQ ID NO:3) and hHGK3 (SEQ ID NO:4) was carried out using the Clustal W program (Thompson, J.D., Higgins, D.G. and Gibson T.J. (1994) *Nuc. Acids. Res.* 22, 4673-4680). Identities are shaded in black and the initiation codons of SGKα (SEQ ID NO:1) and SGKβ (SEQ ID NO:8) are indicated by arrows. The two key phosphorylation sites are marked with asterisks.